



SEQUENCE LISTING

RAPPERSA CORPORATION CALLEN, Walter MATHUR, Eric

<120> ENZYMES HAVING HIGH TEMPERATURE POLYMERASE ACTIVITY AND METHODS OF USE THEREOF

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> US 09/656,309 > 2000-09-06													
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70> PatentIn version 3.0													
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ggt aaa gag cct cag gta atc ata tgg ggt att gct gag aac ggc gag Gly Lys Glu Pro Gln Val Ile Ile Trp Gly Ile Ala Glu Asn Gly Glu 20 25 30	96												
agg gta gtc ctc att gac agg tct ttt cgc cca tac ttc tat gcg ctg Arg Val Val Leu Ile Asp Arg Ser Phe Arg Pro Tyr Phe Tyr Ala Leu 35 40 45	4												
ctt gca ccg ggc gcc gat cct aag cag gta gca caa cgt att cgt gca Leu Ala Pro Gly Ala Asp Pro Lys Gln Val Ala Gln Arg Ile Arg Ala 50 55 60)2												
ttg agt agg cca aag agc ccg att ata ggt gta gag gat gac aag agg Leu Ser Arg Pro Lys Ser Pro Ile Ile Gly Val Glu Asp Asp Lys Arg 65 70 75 80	ł O												
aag tac ttc ggg agg cct cgt agg gtc tta cgt att cgc acc gtg cta Lys Tyr Phe Gly Arg Pro Arg Arg Val Leu Arg Ile Arg Thr Val Leu 85 90 95	8 8												
ccc gag gct gtt agg gag tat cgc gaa ctc gta aag aac gtt gat ggt Pro Glu Ala Val Arg Glu Tyr Arg Glu Leu Val Lys Asn Val Asp Gly 100 105 110	6												
gtt gag gat gtt cta gag gcg gat ata cgc ttc gct atg cgc tat ctc 38	34												

13

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Val	Glu	Asp 115	Val	Leu	Glu	Ala	Asp 120	Ile	Arg	Phe	Ala	Met 125	Arg	Tyr	Leu	
										tac Tyr						432
				_	_			_	_	gac Asp 155	_	_		_	_	480
										gct Ala						528
										att Ile						576
										gta Val						624
										att Ile						672
										tac Tyr 235						720
										cat His						768
										aag Lys						816
					Pro		Thr	Ser	Val	cat His	Gly	His	Val			864
										tat Tyr						912
										gca Ala 315						960
atg Met	aag Lys	aag Lys	agt Ser	gaa Glu 325	cgc Arg	gtt Val	atc Ile	atc Ile	aat Asn 330	tgg Trp	tgg Trp	gag Glu	att Ile	cca Pro 335	gac Asp	1008
										tta Leu						1056

340 345 350

_	_	_	_	_					aag Lys				:	1104
_		_	_	_	_				cta Leu 380	_	_	_	 :	1152
	_	_	_		_	_			ctg Leu		_	_	 :	1200
									gag Glu				:	1248
		_		_	_		-	_	aga Arg				 :	1296
		_	-						cca Pro				:	1344
									cct Pro 460				:	1392
									aag Lys				:	1440
									gag Glu				:	1488
									tat Tyr				:	1536
									gcg Ala				:	1584
									ggc Gly 540				:	1632
									ggt Gly				:	1680
									ctc Leu				<u>:</u>	1728

ggt Gly	gac Asp	aca Thr	gat Asp 580	tcg Ser	ctc Leu	ttc Phe	gtg Val	acc Thr 585	tat Tyr	gat Asp	ccg Pro	gag Glu	aag Lys 590	gtg Val	gaa Glu	1776
aat Asn	ttc Phe	atc Ile 595	aaa Lys	att Ile	ata Ile	aag Lys	gag Glu 600	Glu	ctg Leu	gly aaa	ttc Phe	gaa Glu 605	atc Ile	aag Lys	cta Leu	1824
gag Glu	aag Lys 610	gtg Val	tac Tyr	aaa Lys	cgc Arg	tta Leu 615	ttc Phe	ttt Phe	aca Thr	gag Glu	gct Ala 620	aag Lys	aag Lys	agg Arg	tac Tyr	1872
gct Ala 625	ggc Gly	ctt Leu	ctc Leu	gag Glu	gac Asp 630	gga Gly	cgt Arg	ata Ile	gat Asp	att Ile 635	gtc Val	ggt Gly	ttc Phe	gag Glu	gct Ala 640	1920
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													gta Val 670			2016
gtc Val	agg Arg	aag Lys 675	att Ile	gtg Val	aaa Lys	gag Glu	ttg Leu 680	gag Glu	gag Glu	ggc Gly	aag Lys	gtt Val 685	ccc Pro	ata Ile	gag Glu	2064
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tac Tyr	cgg Arg	gta Val	agc Ser	cca Pro 725	ggc Gly	gac Asp	aag Lys	ata Ile	999 Gly 730	tat Tyr	gta Val	ata Ile	gtg Val	aag Lys 735	ggt Gly	2208
ggt Gly	ggc Gly	cgt Arg	atc Ile 740	agt Ser	caa Gln	aga Arg	gca Ala	tgg Trp 745	cca Pro	tac Tyr	ttc Phe	atg Met	gtc Val 750	aag Lys	gat Asp	2256
cct Pro	agc Ser	cag Gln 755	ata Ile	gac Asp	gtg Val	acc Thr	tac Tyr 760	tat Tyr	gtt Val	gac Asp	cac His	caa Gln 765	atc Ile	atc Ile	ccg Pro	
gct Ala	gca Ala 770	ttg Leu	aga Arg	ata Ile	ctg Leu	ggc Gly 775	tac Tyr	ttt Phe	ggc Gly	atc Ile	acc Thr 780	gag Glu	aag Lys	aag Lys	ctg Leu	2352
aaa Lys 785	gca Ala	agt Ser	gca Ala	act Thr	999 Gly 790	cag Gln	aag Lys	act Thr	ctc Leu	ttc Phe 795	gac Asp	ttt Phe	cta Leu	gcc Ala	aag Lys 800	2400

Y

aag agc aag taa Lys Ser Lys

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<213> Pyrolobus fumarius

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Gly Lys Glu Pro Gln Val Ile Ile Trp Gly Ile Ala Glu Asn Gly Glu 20 25 30

Arg Val Val Leu Ile Asp Arg Ser Phe Arg Pro Tyr Phe Tyr Ala Leu 35 40 45

Leu Ala Pro Gly Ala Asp Pro Lys Gln Val Ala Gln Arg Ile Arg Ala 50 55 60

Leu Ser Arg Pro Lys Ser Pro Ile Ile Gly Val Glu Asp Asp Lys Arg 65 70 75 80

Lys Tyr Phe Gly Arg Pro Arg Arg Val Leu Arg Ile Arg Thr Val Leu 85 90 95

Pro Glu Ala Val Arg Glu Tyr Arg Glu Leu Val Lys Asn Val Asp Gly
100 105 110

Val Glu Asp Val Leu Glu Ala Asp Ile Arg Phe Ala Met Arg Tyr Leu 115 120 125

Ile Asp His Asp Leu Phe Pro Phe Thr Trp Tyr Arg Val Glu Ala Glu 130 135 140

Pro Leu Glu Asn Lys Met Gly Phe Arg Val Asp Lys Val Tyr Leu Val 145 150 155 160

Lys Ser Arg Pro Glu Pro Leu Tyr Gly Glu Ala Leu Ala Pro Thr Lys 165 170 175

Leu Pro Asp Leu Arg Ile Leu Ala Phe Asp Ile Glu Val Tyr Ser Lys

180 185 190

Gln Gly Ser Pro Arg Pro Glu Arg Asp Pro Val Ile Val Ile Ala Val 195 Lys Thr Asp Asp Gly Asp Glu Val Leu Phe Ile Ala Glu Gly Lys Asp Asp Arg Lys Pro Ile Arg Glu Phe Val Glu Tyr Val Lys Arg Tyr Asp Pro Asp Ile Ile Val Gly Tyr Asn Asn His Phe Asp Trp Pro Tyr 250 Leu Leu Arg Arg Ala Arg Ile Leu Gly Ile Lys Leu Asp Val Thr Arg Arg Val Gly Ala Glu Pro Thr Thr Ser Val His Gly His Val Ser Val 280 Pro Gly Arg Leu Asn Val Asp Leu Tyr Asp Tyr Ala Glu Glu Met Pro 295 300 Glu Ile Lys Ile Lys Ser Leu Glu Glu Val Ala Glu Tyr Leu Gly Val 305 310 Met Lys Lys Ser Glu Arg Val Ile Ile Asn Trp Trp Glu Ile Pro Asp 325 Tyr Trp Asp Asp Pro Lys Lys Arg Pro Leu Leu Gln Tyr Ala Arg 345 Asp Asp Val Arg Ala Thr Tyr Gly Leu Ala Glu Lys Ile Leu Pro Phe Ala Ile Gln Leu Ser Tyr Val Thr Gly Leu Pro Leu Asp Gln Val Gly 375 380 Ala Met Ser Val Gly Phe Arg Leu Glu Trp Tyr Leu Ile Arg Ala Ala 385 390 395 Phe Lys Met Lys Glu Leu Val Pro Asn Arg Val Glu Arg Pro Glu Glu

410

415

405

Thr Tyr Arg Gly Ala Ile Val Leu Glu Pro Leu Arg Gly Val His Glu 420 425 430

Asn Ile Ala Val Leu Asp Phe Ser Ser Met Tyr Pro Asn Ile Met Ile 435 440 445

Lys Tyr Asn Val Gly Pro Asp Thr Leu Val Arg Pro Gly Glu Lys Cys 450 455 460

Gly Glu Cys Gly Cys Trp Glu Ala Pro Glu Val Lys His Arg Phe Arg 465 470 475 480

Arg Cys Pro Pro Gly Phe Phe Lys Thr Val Leu Glu Arg Leu Leu Glu 485 490 495

Leu Arg Lys Arg Val Arg Ala Glu Met Lys Lys Tyr Pro Pro Asp Ser 500 505 510

Pro Glu Tyr Arg Leu Leu Asp Glu Arg Gln Lys Ala Leu Lys Val Leu 515 520 525

Ala Asn Ala Ser Tyr Gly Tyr Met Gly Trp Ser Gly Ala Arg Trp Tyr 530 540

Cys Arg Glu Cys Ala Lys Ala Val Thr Ala Trp Gly Arg His Leu Ile 545 550 555 560

Arg Thr Ala Ile Asn Ile Ala Arg Lys Leu Gly Leu Lys Val Ile Tyr 565 570 575

Gly Asp Thr Asp Ser Leu Phe Val Thr Tyr Asp Pro Glu Lys Val Glu 580 585 590

Asn Phe Ile Lys Ile Ile Lys Glu Glu Leu Gly Phe Glu Ile Lys Leu 595 600 605

Glu Lys Val Tyr Lys Arg Leu Phe Phe Thr Glu Ala Lys Lys Arg Tyr 610 615 620

Ala Gly Leu Leu Glu Asp Gly Arg Ile Asp Ile Val Gly Phe Glu Ala 625 630 635 640 Val Arg Gly Asp Trp Cys Glu Leu Ala Lys Glu Val Gln Thr Lys Val 645 650 655

Val Glu Ile Val Leu Lys Thr Ser Glu Val Asn Lys Ala Val Glu Tyr 660 665 670

Val Arg Lys Ile Val Lys Glu Leu Glu Glu Gly Lys Val Pro Ile Glu 675 680 685

Lys Leu Val Ile Trp Lys Thr Leu Ser Lys Arg Leu Glu Glu Tyr Thr 690 695 700

Thr Glu Ala Pro His Val Val Ala Ala Lys Arg Met Leu Ser Ala Gly 705 710 715 720

Tyr Arg Val Ser Pro Gly Asp Lys Ile Gly Tyr Val Ile Val Lys Gly 725 730 735

Gly Gly Arg Ile Ser Gln Arg Ala Trp Pro Tyr Phe Met Val Lys Asp 740 745 750

Pro Ser Gln Ile Asp Val Thr Tyr Tyr Val Asp His Gln Ile Ile Pro 755 760 765

Ala Ala Leu Arg Ile Leu Gly Tyr Phe Gly Ile Thr Glu Lys Lys Leu 770 775 780

Lys Ala Ser Ala Thr Gly Gln Lys Thr Leu Phe Asp Phe Leu Ala Lys 785 790 795 800

Lys Ser Lys